

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/972,758A

DATE: 04/18/2003 TIME: 15:54:26

Input Set : A:\277084004.ST25.txt

Output Set: N:\CRF4\04182003\I972758A.raw

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3 <110> APPLICANT: Case Western Reserve University
         Montano, Monica
         Wittman, Bryan
 7 <120> TITLE OF INVENTION: Suppressors of Human Breast Cancer Cell Growth
 9 <130> FILE REFERENCE: 27708/04004
11 <140> CURRENT APPLICATION NUMBER: US 09/972758A
12 <141> CURRENT FILING DATE: 2001-10-05
14 <150> PRIOR APPLICATION NUMBER: US 60/238,187
15,<151> PRIOR FILING DATE: 2000-10-05
17 <160> NUMBER OF SEQ ID NOS: 7
                                                         ENTERED
19 <170> SOFTWARE: PatentIn version 3.1
21 <210> SEQ ID NO: 1
22 <211> LENGTH: 1080
23 <212> TYPE: DNA
24 <213> ORGANISM: Homo sapiens
26 <400> SEQUENCE: 1
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29 gctgctgctg tccaggaaga gctgaaccct gagcgccccc caggcgcgga ggagcgggtg
                                                                        120
31 cccgaggagg acagtaggtg gcaatcgaga gcgttccccc agttgggtgg ccgtccgggg
                                                                        180
33 ccggagggg aagggagcct ggaatcccaa ccacctccct tgcagaccca ggcctgtcca
                                                                        240
35 gaatetaget geetgagaga gggegagaag ggeeagaatg gggaegaete gteegetgge
                                                                        300
37 ggcgaettee egeegeegge agaagtggaa eegaegeeeg aggeegaget getegeeeag
                                                                        360
39 cettgteatg acteegagge eagtaagttg ggggeteetg eegeaggggg egaagaggag
                                                                        420
41 tggggacagc agcagagaca gctggggaag aaaaaacata agagacgccc gtccaagaag
                                                                        480
43 aagcggcatt ggaaaccgta ctacaagctg aactgggaag agaagaaaaa gttcgacgag
                                                                        540
45 aaacagagee ttegagette aaggateega geegagatgt tegeeaaggg eeageeggte
                                                                        600
47 gcgccctata acaccacgca gttcctcatg gatgatcacg accaggagga gccggatctc
                                                                        660
49 aaaaccggcc tgtactccaa gcgggccgcc gccaaatccg acgacaccag cgatgacgac
                                                                        720
51 ttcatggaag aagggggtga ggaggatggg ggcagcgatg ggatgggagg ggacggcagc
                                                                        780
53 gagtttctgc agcgggactt ctcggagacg tacgagcggt accacacgga gagcctgcag
                                                                        840
55 aacatgagca agcaggagct catcaaggag tacctggaac tggagaagtg cctctcgcgc
                                                                        900
57 atggaggacg agaacaaccg gctgcggctg gagagcaagc ggctgggtgg cgacgacgcg
                                                                        960
59 cgtgtgcggg agctggagct ggagctggac cggctgcgcg ccgagaacct ccagctgctg
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65 <211> LENGTH: 359
66 <212> TYPE: PRT
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71 Met Ala Glu Pro Phe Leu Ser Glu Tyr Gln His Gln Pro Gln Thr Ser
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75 Asn Cys Thr Gly Ala Ala Ala Val Gln Glu Glu Leu Asn Pro Glu Arg
76
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79 Pro Pro Gly Ala Glu Glu Arg Val Pro Glu Glu Asp Ser Arg Trp Gln
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 83 Ser Arg Ala Phe Pro Gln Leu Gly Gly Arg Pro Gly Pro Glu Gly Glu
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" 87 Gly Ser Leu Glu Ser Gln Pro Pro Pro Leu Gln Thr Gln Ala Cys Pro
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                                           75
 91 Glu Ser Ser Cys Leu Arg Glu Gly Glu Lys Gly Gln Asn Gly Asp Asp
                                        90
 95 Ser Ser Ala Gly Gly Asp Phe Pro Pro Pro Ala Glu Val Glu Pro Thr
                100
                                    105
 99 Pro Glu Ala Glu Leu Leu Ala Gln Pro Cys His Asp Ser Glu Ala Ser
 100 115
                                 120
 103 Lys Leu Gly Ala Pro Ala Ala Gly Gly Glu Glu Glu Trp Gly Gln Gln
                            135
 107 Gln Arg Gln Leu Gly Lys Lys His Arg Arg Arg Pro Ser Lys Lys
                         150
                                            155
 111 Lys Arg His Trp Lys Pro Tyr Tyr Lys Leu Thr Trp Glu Glu Lys Lys
                    165
                                        170
 115 Lys Phe Asp Glu Lys Gln Ser Leu Arg Ala Ser Arg Ile Arg Ala Glu
                180
                                    185
 119 Met Phe Ala Lys Gly Gln Pro Val Ala Pro Tyr Asn Thr Thr Gln Phe
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 123 Leu Met Asp Asp His Asp Gln Glu Glu Pro Asp Leu Lys Thr Gly Leu
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 127 Tyr Ser Lys Arg Ala Ala Ala Lys Ser Asp Asp Thr Ser Asp Asp
                         230
                                            235
 131 Phe Met Glu Glu Gly Gly Glu Glu Asp Gly Gly Ser Asp Gly Met Gly
                     245
                                       250
 135 Gly Asp Gly Ser Glu Phe Leu Gln Arg Asp Phe Ser Glu Thr Tyr Glu
 136 .
                260
                                    265
 139 Arg Tyr His Thr Glu Ser Leu Gln Asn Met Ser Lys Gln Glu Leu Ile
           275
                                280
 143 Lys'Glu Tyr Leu Glu Leu Glu Lys Cys Leu Ser Arg Met Glu Asp Glu
                            295
 147. Asn Asn Arg Leu Arg Leu Glu Ser Lys Arg Leu Gly Gly Asp Asp Ala
                        310
                                            315
 151 Arg Val Arg Glu Leu Glu Leu Glu Leu Asp Arg Leu Arg Ala Glu Asn
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 155 Leu Gln Leu Leu Thr Glu Asn Glu Leu His Arg Gln Gln Glu Arg Ala
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 159 Pro Leu Ser Lys Phe Gly Asp
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 164 <211> LENGTH: 19
 165 <212> TYPE: PRT
 166 <213> ORGANISM: Homo sapiens
 168 <400> SEQUENCE: 3
 170 Lys His Arg Arg Pro Ser Lys Lys Arg His Trp Lys Pro Tyr
 171 1
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	208	<213> ORGANISM: Homo sapiens	
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VERIFICATION SUMMARY

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